

## **RAW SEQUENCE LISTING**

**The Biotechnology Systems Branch of the Scientific and Technical  
Information Center (STIC) no errors detected.**

Application Serial Number: 10/573,962  
Source: IFWP  
Date Processed by STIC: 4/7/06

# ***ENTERED***



IFWP

## RAW SEQUENCE LISTING

DATE: 04/07/2006

PATENT APPLICATION: US/10/573,962

TIME: 14:58:51

Input Set : A:\PTO.DA.txt

Output Set: N:\CRF4\04072006\J573962.raw

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3 <110> APPLICANT: GOVERNMENT OF THE UNITED STATES OF AMERICA, REPRESENTED BY
4     THE SECRETARY, DEPARTMENT OF HEALTH AND HUMAN SERVICES
5     DIMITROV, Dimiter S.
6     ZHANG, Mei-Yun
8 <120> TITLE OF INVENTION: IMMUNOGLOBULINS WITH POTENT AND BROAD ANTIVIRAL ACTIVITY
10 <130> FILE REFERENCE: 251149
C--> 12 <140> CURRENT APPLICATION NUMBER: US/10/573,962
C--> 12 <141> CURRENT FILING DATE: 2006-03-29
12 <150> PRIOR APPLICATION NUMBER: US 60/506,946
13 <151> PRIOR FILING DATE: 2003-09-29
15 <150> PRIOR APPLICATION NUMBER: PCT/US04/31878
16 <151> PRIOR FILING DATE: 2004-09-29
18 <160> NUMBER OF SEQ ID NOS: 19
20 <170> SOFTWARE: PatentIn version 3.3
22 <210> SEQ ID NO: 1
23 <211> LENGTH: 254
24 <212> TYPE: PRT
25 <213> ORGANISM: Artificial
27 <220> FEATURE:
28 <223> OTHER INFORMATION: Synthetic
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33 1          5          10          15
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37          20          25          30
40 Ala Trp Tyr Gln Gln Lys Pro Gly Gln Ala Pro Arg Leu Leu Ile Tyr
41          35          40          45
44 Gly Ala Ser Thr Arg Ala Thr Gly Ile Pro Asp Arg Phe Ser Gly Ser
45          50          55          60
48 Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Gly Arg Leu Glu Pro Glu
49 65          70          75          80
52 Asp Leu Ala Val Tyr Tyr Cys Gln Gln Tyr Gly Thr Ser Pro Tyr Thr
53          85          90          95
56 Phe Gly Gln Gly Thr Lys Leu Glu Ile Lys Arg Thr Gly Gly Gly Gly
57          100         105         110
60 Ser Gly Gly Gly Ala Ser Gly Gly Gly Ser Val Arg Leu Leu Glu
61          115         120         125
64 Gln Ser Gly Ala Glu Val Lys Lys Pro Gly Ser Ser Val Gln Val Ser
65          130         135         140
68 Cys Lys Ala Ser Gly Gly Thr Phe Ser Met Tyr Gly Val Asn Trp Val
69 145         150         155         160
72 Arg Gln Ala Pro Gly His Gly Leu Glu Trp Met Gly Gly Ile Ile Pro
73          165         170         175

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76 Ile Phe Gly Thr Ser Asn Tyr Ala Gln Lys Phe Arg Gly Arg Val Thr
77           180           185           190
80 Phe Thr Ala Asp Gln Ala Thr Ser Thr Ala Tyr Met Glu Leu Thr Asn
81           195           200           205
84 Leu Arg Ser Asp Asp Thr Ala Val Tyr Tyr Cys Ala Arg Asp Phe Gly
85           210           215           220
88 Pro Asp Trp Glu Asp Gly Asp Ser Tyr Asp Gly Ser Gly Arg Gly Phe
89 225           230           235           240
92 Phe Asp Phe Trp Gly Gln Gly Thr Leu Val Thr Val Ser Ser
93           245           250
96 <210> SEQ ID NO: 2
97 <211> LENGTH: 254
98 <212> TYPE: PRT
99 <213> ORGANISM: Artificial
101 <220> FEATURE:
102 <223> OTHER INFORMATION: Synthetic
104 <400> SEQUENCE: 2
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107 1           5           10           15
110 Ala Thr Leu Ser Cys Arg Ala Ser Gln Ser Val Ser Ser Gly Ser Leu
111           20           25           30
114 Ala Trp Tyr Gln Gln Lys Pro Gly Gln Ala Pro Arg Leu Leu Ile Tyr
115           35           40           45
118 Gly Ala Ser Thr Arg Ala Thr Gly Ile Pro Asp Arg Phe Ser Gly Ser
119           50           55           60
122 Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Gly Arg Leu Glu Pro Glu
123 65           70           75           80
126 Asp Leu Ala Val Tyr Tyr Cys Gln Gln Tyr Gly Thr Ser Pro Tyr Thr
127           85           90           95
130 Phe Gly Gln Gly Thr Lys Leu Glu Ile Lys Arg Thr Gly Gly Gly Gly
131           100          105          110
134 Ser Gly Gly Gly Gly Ser Gly Gly Gly Gly Ser Val Gln Leu Leu Glu
135           115          120          125
138 Gln Ser Gly Ala Glu Val Lys Lys Pro Gly Ser Ser Val Gln Val Ser
139           130          135          140
142 Cys Lys Ala Phe Gly Gly Thr Phe Ser Met Tyr Gly Phe Asn Trp Val
143 145           150           155           160
146 Arg Gln Ala Pro Gly His Gly Leu Glu Trp Met Gly Gly Ile Ile Pro
147           165          170          175
150 Ile Phe Gly Thr Thr Asn Tyr Ala Gln Lys Phe Arg Gly Arg Val Thr
151           180          185          190
154 Phe Thr Ala Asp Gln Ala Thr Ser Thr Ala Tyr Met Glu Leu Thr Asn
155           195          200          205
158 Leu Arg Ser Asp Asp Thr Ala Val Tyr Tyr Cys Ala Arg Asp Phe Gly
159           210          215          220
162 Pro Asp Trp Glu Gly Gly Asp Ser Tyr Asp Gly Ser Gly Arg Gly Phe
163 225           230          235           240
166 Phe Asp Phe Trp Gly Gln Gly Thr Leu Val Asn Val Ser Ser
167           245          250

```

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170 <210> SEQ ID NO: 3
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172 <212> TYPE: PRT
173 <213> ORGANISM: Artificial
175 <220> FEATURE:
176 <223> OTHER INFORMATION: Synthetic
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188 <210> SEQ ID NO: 4
189 <211> LENGTH: 34
190 <212> TYPE: PRT
191 <213> ORGANISM: Artificial
193 <220> FEATURE:
194 <223> OTHER INFORMATION: Synthetic
196 <400> SEQUENCE: 4
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199 1 5 10 15
202 Asp Pro Glu Glu Pro Lys Ser Cys Asp Lys Thr His Thr Cys Pro Pro
203 20 25 30
206 Cys Pro
210 <210> SEQ ID NO: 5
211 <211> LENGTH: 216
212 <212> TYPE: PRT
213 <213> ORGANISM: Artificial
215 <220> FEATURE:
216 <223> OTHER INFORMATION: Synthetic
218 <400> SEQUENCE: 5
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221 1 5 10 15
224 Pro Lys Asp Thr Leu Met Ile Ser Arg Thr Pro Glu Val Thr Cys Val
225 20 25 30
228 Val Val Asp Val Ser His Glu Asp Pro Glu Val Lys Phe Asn Trp Tyr
229 35 40 45
232 Val Asp Gly Val Glu Val His Asn Ala Lys Thr Lys Pro Arg Glu Glu
233 50 55 60
236 Gln Tyr Asn Ser Thr Tyr Arg Val Val Ser Val Leu Thr Val Leu His
237 65 70 75 80
240 Gln Asp Trp Leu Asn Gly Lys Glu Tyr Lys Cys Lys Val Ser Asn Lys
241 85 90 95
244 Ala Leu Pro Ala Pro Ile Glu Lys Thr Ile Ser Lys Ala Lys Gly Gln
245 100 105 110
248 Pro Arg Glu Pro Gln Val Tyr Thr Leu Pro Pro Ser Arg Asp Glu Leu
249 115 120 125
252 Thr Lys Asn Gln Val Ser Leu Thr Cys Leu Val Lys Gly Phe Tyr Pro
253 130 135 140
256 Ser Asp Ile Ala Val Glu Trp Glu Ser Asn Gly Gln Pro Glu Asn Asn
257 145 150 155 160

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260 Tyr Lys Thr Thr Pro Pro Val Leu Asp Ser Asp Gly Ser Phe Phe Leu
261                               165                               170                               175
264 Tyr Ser Lys Leu Thr Val Asp Lys Ser Arg Trp Gln Gln Gly Asn Val
265                               180                               185                               190
268 Phe Ser Cys Ser Val Met His Glu Ala Leu His Asn His Tyr Thr Gln
269                               195                               200                               205
272 Lys Ser Leu Ser Leu Ser Pro Gly
273       210                               215
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277 <211> LENGTH: 489
278 <212> TYPE: PRT
279 <213> ORGANISM: Artificial
281 <220> FEATURE:
282 <223> OTHER INFORMATION: Synthetic
284 <400> SEQUENCE: 6
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287 1                               5                               10                               15
290 Ala Thr Leu Ser Cys Arg Ala Ser Gln Ser Val Ser Ser Gly Ser Leu
291                               20                               25                               30
294 Ala Trp Tyr Gln Gln Lys Pro Gly Gln Ala Pro Arg Leu Leu Ile Tyr
295                               35                               40                               45
298 Gly Ala Ser Thr Arg Ala Thr Gly Ile Pro Asp Arg Phe Ser Gly Ser
299       50                               55                               60
302 Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Gly Arg Leu Glu Pro Glu
303 65                               70                               75                               80
306 Asp Leu Ala Val Tyr Tyr Cys Gln Gln Tyr Gly Thr Ser Pro Tyr Thr
307                               85                               90                               95
310 Phe Gly Gln Gly Thr Lys Leu Glu Ile Lys Arg Thr Gly Gly Gly Gly
311                               100                              105                              110
314 Ser Gly Gly Gly Ala Ser Gly Gly Gly Gly Ser Val Arg Leu Leu Glu
315                               115                              120                              125
318 Gln Ser Gly Ala Glu Val Lys Lys Pro Gly Ser Ser Val Gln Val Ser
319       130                              135                              140
322 Cys Lys Ala Ser Gly Gly Thr Phe Ser Met Tyr Gly Val Asn Trp Val
323 145                              150                              155                              160
326 Arg Gln Ala Pro Gly His Gly Leu Glu Trp Met Gly Gly Ile Ile Pro
327                               165                               170                               175
330 Ile Phe Gly Thr Ser Asn Tyr Ala Gln Lys Phe Arg Gly Arg Val Thr
331                               180                               185                               190
334 Phe Thr Ala Asp Gln Ala Thr Ser Thr Ala Tyr Met Glu Leu Thr Asn
335       195                               200                               205
338 Leu Arg Ser Asp Asp Thr Ala Val Tyr Tyr Cys Ala Arg Asp Phe Gly
339       210                               215                               220
342 Pro Asp Trp Glu Asp Gly Asp Ser Tyr Asp Gly Ser Gly Arg Gly Phe
343 225                               230                               235                               240
346 Phe Asp Phe Trp Gly Gln Gly Thr Leu Val Thr Val Ser Ser Pro Asp
347                               245                               250                               255
350 Pro Glu Glu Pro Lys Ser Cys Asp Lys Thr His Thr Cys Pro Pro Cys
351       260                               265                               270

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354 Pro Ala Pro Glu Leu Leu Gly Gly Pro Ser Val Phe Leu Phe Pro Pro
355      275      280      285
358 Lys Pro Lys Asp Thr Leu Met Ile Ser Arg Thr Pro Glu Val Thr Cys
359      290      295      300
362 Val Val Val Asp Val Ser His Glu Asp Pro Glu Val Lys Phe Asn Trp
363 305      310      315      320
366 Tyr Val Asp Gly Val Glu Val His Asn Ala Lys Thr Lys Pro Arg Glu
367      325      330      335
370 Glu Gln Tyr Asn Ser Thr Tyr Arg Val Val Ser Val Leu Thr Val Leu
371      340      345      350
374 His Gln Asp Trp Leu Asn Gly Lys Glu Tyr Lys Cys Lys Val Ser Asn
375      355      360      365
378 Lys Ala Leu Pro Ala Pro Ile Glu Lys Thr Ile Ser Lys Ala Lys Gly
379      370      375      380
382 Gln Pro Arg Glu Pro Gln Val Tyr Thr Leu Pro Pro Ser Arg Asp Glu
383 385      390      395      400
386 Leu Thr Lys Asn Gln Val Ser Leu Thr Cys Leu Val Lys Gly Phe Tyr
387      405      410      415
390 Pro Ser Asp Ile Ala Val Glu Trp Glu Ser Asn Gly Gln Pro Glu Asn
391      420      425      430
394 Asn Tyr Lys Thr Thr Pro Pro Val Leu Asp Ser Asp Gly Ser Phe Phe
395      435      440      445
398 Leu Tyr Ser Lys Leu Thr Val Asp Lys Ser Arg Trp Gln Gln Gly Asn
399      450      455      460
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407      485
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411 <211> LENGTH: 504
412 <212> TYPE: PRT
413 <213> ORGANISM: Artificial
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416 <223> OTHER INFORMATION: Synthetic
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425      20      25      30
428 Ala Trp Tyr Gln Gln Lys Pro Gly Gln Ala Pro Arg Leu Leu Ile Tyr
429      35      40      45
432 Gly Ala Ser Thr Arg Ala Thr Gly Ile Pro Asp Arg Phe Ser Gly Ser
433      50      55      60
436 Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Gly Arg Leu Glu Pro Glu
437 65      70      75      80
440 Asp Leu Ala Val Tyr Tyr Cys Gln Gln Tyr Gly Thr Ser Pro Tyr Thr
441      85      90      95
444 Phe Gly Gln Gly Thr Lys Leu Glu Ile Lys Arg Thr Gly Gly Gly Gly
445      100      105      110

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RAW SEQUENCE LISTING ERROR SUMMARY  
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Input Set : A:\PTO.DA.txt  
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Invalid <213> Response:

Use of "Artificial" only as "<213> Organism" response is incomplete,  
per 1.823(b) of New Sequence Rules. Valid response is Artificial Sequence.

Seq#:1,2,3,4,5,6,7,8,9,10,11,12,13,14,15,16,17,18,19

**VERIFICATION SUMMARY**

DATE: 04/07/2006

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Input Set : A:\PTO.DA.txt

Output Set: N:\CRF4\04072006\J573962.raw

L:12 M:270 C: Current Application Number differs, Replaced Current Application No

L:12 M:271 C: Current Filing Date differs, Replaced Current Filing Date